

REFERENCE
AUTHORS

2 (bases 1 to 154902)
 Birren, B., Linton, J., Nusbaum, C., Lander, E., Allon, N., Anderson, M., Baker, J., Baldwin, J., Barna, N., Beckwith, J., Bernier, J., Bourgeois, C., Brown, A., Castle, A., Cerny, J., Collange, M., Collins, S., Collymore, A., Cooke, P., Corliss, D., Deyayre, R., Denon, K., Dewar, K., Doncelan, L., Ferreira, P., FitzHugh, W., Forrest, C., Fuxer, R., Gage, E., Gardyna, S., Geraghty, K., Grant, G., Hades, B., Heaford, A., Hendon, A., Horton, L., Huxland, J., Jancot, L., Jones, C., Kann, L., Karas, A., Lebecky, J., Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K., Melnick, J., Molla, M., Morris, W., Morrow, J., Muchaleckyj, J., Naf, K., Naylor, J., Nilsson, M., O'Connor, T., Pavlin, P., Pavlin, P., Peterson, K., Riley, R., Roberts, D., Roy, A., Severy, P., Stange-Thomann, N., Stillwell, J., Stojanovic, N., Stone, C., Subramanian, A., Tesfaye, S., Tichovolsky, N., Tortorella-Miller, J., Vassiliev, H., Vo, A., Wagner, A., Wheeler, J., Wu, Y., Wyman, D., Ye, W., Zhao, J., and Zody, M.

TITLE
JOURNAL

Submitted (06-OCT-1998) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA
 3 (bases 1 to 154902)

REFERENCE
AUTHORS

Birren, B., Linton, J., Nusbaum, C., Lander, E., Allon, N., Anderson, M., Baker, J., Baldwin, J., Barna, N., Beckwith, J., Bernier, J., Bourgeois, C., Brown, A., Castle, A., Cerny, J., Collange, M., Collins, S., Collymore, A., Cooke, P., Corliss, D., Deyayre, R., Denon, K., Dewar, K., Doncelan, L., Ferreira, P., FitzHugh, W., Forrest, C., Fuxer, R., Gage, E., Gardyna, S., Geraghty, K., Grant, G., Hades, B., Heaford, A., Hendon, A., Horton, L., Huxland, J., Jancot, L., Jones, C., Kann, L., Karas, A., Lebecky, J., Macdonald, P., Marquis, N., McEwan, P., McGurk, A., McKernan, K., Melnick, J., Molla, M., Morris, W., Morrow, J., Muchaleckyj, J., Naf, K., Naylor, J., Nilsson, M., O'Connor, T., Pavlin, P., Pavlin, P., Peterson, K., Riley, R., Roberts, D., Roy, A., Severy, P., Stange-Thomann, N., Stillwell, J., Stojanovic, N., Stone, C., Subramanian, A., Tesfaye, S., Tichovolsky, N., Tortorella-Miller, J., Vassiliev, H., Vo, A., Wagner, A., Wheeler, J., Wu, Y., Wyman, D., Ye, W., Zhao, J., and Zody, M.

TITLE
JOURNAL

Submitted (24-OCT-1998) Whitehead Institute/MIT Center for Genome Research, 320 Charles Street, Cambridge, MA 02141, USA
 On Oct 24, 1998 this sequence version replaced gi:3766128.

COMMENT

All repeats were identified using RepeatMasker: Smit, A.F.A. & Green, P. (1996-1997)

FEATURES

source

location/Ceallifiers

1. 154902

/organism="Homo sapiens"

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/map="17"

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3196..3286

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4435..4746

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5591..5695

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complement(7321..7530)

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9643..9672

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10024..10056

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11327..11373

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complement(11545..11584)

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complement(11599..11652)

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complement(13873..13920)

/note="Stranded coverage."

complement(16795..16816)

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complement(19373..19411)

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19414..19543

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complement(23621..23673)

/rpt_family="MER5A"

complement(27233..27314)

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28133..28224

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complement(29411..29433)

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29378..29379

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35564..35782

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E0 508 CAGGTA 515
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RESULT 7
AK097540
LOCUS
DEFINITION Homo sapiens cDNA FLJ40261 f15, clone TEST12055679, highly similar to CAMP-DEPENDENT PROTEIN KINASE TYPE 1 ALPHA REGULATORY CHAIN.
ACCESSION AK097540
VERSION 1
KEYWORDS 5' UTR capping; f15 (full insert sequence);
SOURCE Homo sapiens testis cDNA to mPNA, clone LIB:TEST12
      CDS:TEST12055679.

ORGANISM Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominoidea; Homo;
1
AUTHORS Kawakami,H., Sugiyama,A., Takekoshi,M., Sugiyama,T., Iida,E.,
Otsuki,T., Sato,H., Wakamatsu,A., Ishii,S., Yamamoto,J., Isono,Y.,
Kawai-Hio,Y., Saito,K., Nishikawa,T., Kimura,K., Yamashita,H.,
Matsuo,K., Nakamura,Y., Sekize,M., Kikuchi,H., Kanda,K.,
Wagatsuna,M., Murakawa,K., Kanchori,K., Takahashi-Fujii,A.,
Oshima,A., Suzuki,Y., Sujiato,S., Naqurari,K., Masubo,Y., Nagai,K.
and Tsouai,T.
TITLE NEDC human cDNA sequencing project.
JOURNAL Unpublished
REFERENCE 2 (bases 1 to 1940)
AUTHORS Tsouai,T. and Yamamoto,J.
TITLE Direct Submission
JOURNAL Submitted (04-JUL-2002) Takao Tsouai, FLJ Project (HRI Team); 2-6-7
Kazusa-Kametari, Kisarazu, Chiba 292-0812, Japan
(-mail:genomics@kazusa.or.jp, Tel:81-448-52-3975, Fax:81-448-52-3386)
NEDC human cDNA sequencing project supported by Ministry of
Economy, Trade and Industry of Japan; cDNA full insert sequencing:
Research Association for Biotechnology (RAB); cDNA library
construction: Helix Research Institute (HRI) (supported by Japan
Key Technology Center etc.); 5' & 3'-end one pass sequencing: RAB,
HRI, and Biotechnology Center, National Institute of Technology and
Evaluation; clone selection for full insert sequencing: HRI and
RAB; annotation: HRI and RAB.
FEATURES
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1a in GenBank Accession Number U06412"
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